The following <u>Listing of the Claims</u> will replace all prior versions and all prior listings of the claims in the present application:

<u>Listing of the Claims</u>

56.   (Currently Amended) A method for screening a first repertoire of members comprising a heavy or light chain polypeptide against a second repertoire of members comprising a heavy or light chain polypeptide to identify those members of the first repertoire which interact with members of the second repertoire, comprising :

(a)   arranging the first <u>repertoire in at least one first series of continuous lines wherein each line of said first series comprises a member of said first repertoire</u> and <u>arranging the</u> second repertoires in at least ~~two~~ <u>one second</u> series of continuous lines <u>wherein each line of said second series comprises a member of said second repertoire,</u> ~~to~~ <u>wherein the first and second repertoires</u> form an array, and wherein a plurality of members of the first repertoire are juxtaposed to a plurality of members of the second repertoire; and

(b)   ~~detecting an interaction between heavy or light chain polypeptides of the first and~~ second repertoires, thereby identifying those members of the first repertoire that interact with members of the second repertoire.

57.   (Previously Added) The method of claim 56, wherein said first and second repertoires are each present in a series of continuous, non-intersecting lines.

58.   (Previously Added) The method of claim 56, wherein said heavy or light chain polypeptide is a domain antibody (dAb).

59.   (Previously Added) The method of claim 56, wherein said first repertoire comprises V_H or V_L.

60.   (Previously Added) The method of claim 56, wherein said second repertoire comprises V_H or V_L.